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TECH CENTER 1600/2000

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SEQUENCE LISTING

<110> Anderson, Christen M.
Davis, Robert E.
Clevenger, William
Wiley, Sandra Eileen
Willer, Scott W.
Szabo, Tomas R.
Ghosh, Soumitra S.

<120> PRODUCTION OF ADENINE NUCLEOTIDE
TRANSLOCATOR (ANT), NOVEL ANT LIGANDS AND SCREENING ASSAYS
THEREFOR

<130> 660088.420

<140> US 09/185,904

<141> 1999-11-17

<160> 33

<170> FastSEQ for Windows Version 3.0

<210> 1

<211> 541

<212> DNA

<213> Homo sapien

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caagtgagc	gggtgtgtc	ctacacctt	gacaatttc	gtctatgat	gatgatgag	720
tgggtggga	aaaggggga	tattatgtc	aggggagag	ttactatg	aggaagagt	780
caagtgagc	agggagcaa	gggtctctt	caagtgagc	gggtctctt	ctctatgat	840
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tacttccaa	ccaaagctt	ccctctctt	ttaaggaaa	agtaaaaca	ccctctctt	300

ggtaggtgtg	acaagagaac	ccagttttgg	egetaetttg	cagagaatct	ggcatcgggt	360
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<213> Homo sapien

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gcacacaaac	agttcttggc	cgacacagag	tacaagggtt	tgtgtgattc	catcttctcc	180
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taattcccca	cttaagggct	caattctctc	ttcaagggtt	agttcaacaa	taattctctc	300
gtgtgttgg	cttaagggct	caattctctc	aggtactttg	caggacacat	ggtctctctc	360
gtgtgttgg	cttaagggct	caattctctc	gtgtactttg	tggtatttgc	caggacacac	420
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cagtcagggc	gcaaaaggag	tgcacatcat	tacaagggca	cgttcgactg	ttggagggaac	780
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<220>

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 <220>
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 <210> 7
 <211> 43
 <212> DNA
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 <220>
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 <400> 7
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 <220>
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 <400> 8
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 <211> 44
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> PCR primer

 <400> 9
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 <210> 10
 <211> 44
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Sequencing primer

 <400> 10
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 <210> 11
 <211> 45
 <212> DNA
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<220>		
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cgccaaaaca gccaaact	18	
<210> 12		
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taactgtacg accatgaaga taagatgacg gaacaggaca tctcc	45	
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<223> PCR primer		
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<210> 17
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<220>
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<400> 17
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<210> 18
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<400> 18
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<400> 19
 ccttcacaca gattttc 48

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 atttcgatt cccatata 48

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<210> 23
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<211> 41

<212> ENA

<213> Artificial Sequence

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41

<210> 28

<211> 42

<212> ENA

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<220>

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42

<210> 29

<211> 42

<212> ENA

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<211> 15

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<220>

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<210> 31

<211> 297

<212> PRT

<213> Homo sapiens

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Val Ala Ala Ala Val Ser Lys Thr Ala Val Ala Phe Ile Thr Arg Val
40

Lys Leu Leu Leu Gln Val Gln His Ala Ser Lys Gln Ile Ser Ala Glu
 35 40 45
 Lys Gln Tyr Lys Gly Ile Ile Asp Cys Val Val Arg Ile Pro Lys Glu
 50 55 60
 Gln Gly Phe Leu Ser Phe Trp Arg Gly Asn Leu Ala Asn Val Ile Arg
 65 70 75 80
 Tyr Phe Pro Thr Gln Ala Leu Asn Phe Ala Phe Lys Asp Lys Tyr Lys
 85 90 95
 Gln Leu Phe Leu Gly Gly Val Asp Arg His Lys Gln Phe Trp Arg Tyr
 100 105 110
 Phe Ala Gly Asn Leu Ala Ser Gly Gly Ala Ala Gly Ala Thr Ser Leu
 115 120 125
 Cys Phe Val Tyr Pro Leu Asp Phe Ala Arg Thr Arg Leu Ala Ala Asp
 130 135 140
 Val Gly Arg Arg Ala Gln Arg Glu Phe His Gly Leu Gly Asp Cys Ile
 145 150 155 160
 Ile Lys Ile Phe Lys Ser Asp Gly Leu Arg Gly Leu Tyr Gln Gly Phe
 165 170 175
 Asn Val Ser Val Gln Gly Ile Ile Ile Tyr Arg Ala Ala Tyr Phe Gly
 180 185 190
 Val Tyr Asp Thr Ala Lys Gly Met Leu Pro Asp Pro Lys Asn Val His
 195 200 205
 Ile Phe Val Ser Trp Met Ile Ala Gln Ser Val Thr Ala Val Ala Gly
 210 215 220
 Leu Leu Ser Tyr Pro Phe Asp Thr Val Ala Arg Arg Met Met Met Gln
 225 230 235 240
 Ser Gly Arg Lys Gly Ala Asp Ile Met Tyr Thr Gly Thr Val Asp Cys
 245 250 255
 Trp Arg Lys Ile Ala Lys Asp Glu Gly Ala Lys Ala Phe Phe Lys Gly
 260 265 270
 Ala Trp Ser Asn Val Leu Arg Gly Met Gly Gly Ala Phe Val Leu Val
 275 280 285
 Leu Tyr Asp Glu Ile Lys Lys Tyr Val
 290 295

<210> 32

<211> 238

<212> PRT

<213> Homo sapien

<400> 32

Met Thr Asp Ala Ala Leu Ser Ile Ala Lys Asp Phe Leu Ala Gly Gly
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 15 20 25
 Lys Leu Leu Leu Gln Val Gln His Ala Ser Lys Gln Ile Thr Ala Arg
 30 35 40 45
 Lys Gln Tyr Lys Gly Ile Ile Asp Cys Val Val Arg Ile Pro Lys Glu
 50 55 60
 Gln Gln Val Leu Ser Ile Trp Arg Gly Asn Leu Ala Asn Val Ile Arg
 65 70 75 80
 Tyr Phe Pro Thr Gln Ala Leu Asn Ile Ala Ile Lys Asp Lys Tyr Lys
 85 90 95
 Gln Ile Phe Leu Gly Gly Val Asp Lys Arg Thr Gln Phe Trp Arg Tyr
 100 105 110
 Phe Ala Gly Asn Leu Ala Ser Gly Gly Ala Ala Gly Ala Thr Ser Leu
 115 120 125

Cys Phe Val Tyr Pro Leu Asp Phe Ala Arg Thr Arg Leu Ala Ala Asp
 130 135 140
 Val Gly Lys Ala Gly Ala Glu Arg Glu Phe Arg Gly Leu Gly Asp Cys
 145 150 155 160
 Leu Val Lys Ile Tyr Lys Ser Asp Gly Ile Lys Gly Leu Tyr Gln Gly
 165 170 175
 Phe Asn Val Ser Val Gln Gly Ile Ile Ile Tyr Arg Ala Ala Tyr Phe
 180 185 190
 Gly Ile Tyr Asp Thr Ala Lys Gly Met Leu Pro Asp Pro Lys Asn Thr
 195 200 205
 His Ile Val Ile Ser Trp Met Ile Ala Gln Thr Val Thr Ala Val Ala
 210 215 220
 Gly Leu Thr Ser Tyr Pro Phe Asp Thr Val Arg Arg Arg Met Met Met
 225 230 235 240
 Gln Ser Gly Arg Lys Gly Thr Asp Ile Met Tyr Thr Gly Thr Leu Asp
 245 250 255
 Cys Trp Arg Lys Ile Ala Arg Asp Gln Gly Gly Lys Ala Phe Phe Lys
 260 265 270
 Gly Ala Trp Ser Asn Val Leu Arg Gly Met Gly Gly Ala Phe Val Leu
 275 280 285
 Val Leu Tyr Asp Glu Ile Lys Lys Tyr Thr
 290 295

2110- 44

2111- 298

2112- FRT

2113- Home sapien

2400- 33

Met Thr Glu Gln Ala Ile Ser Phe Ala Lys Asp Phe Leu Ala Gly Gly
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 Ile Ala Ala Ala Ile Ser Lys Thr Ala Val Ala Pro Ile Glu Arg Val
 20 25 30
 Lys Leu Leu Leu Gln Val Gln His Ala Ser Lys Gln Ile Ala Ala Asp
 35 40 45
 Lys Gln Tyr Lys Gly Ile Val Asp Cys Ile Val Arg Ile Pro Lys Glu
 50 55 60
 Gln Gly Val Leu Ser Phe Trp Arg Gly Asn Leu Ala Asn Val Ile Arg
 65 70 75 80
 Tyr Phe Pro Thr Gln Ala Leu Asn Phe Ala Phe Lys Asp Lys Tyr Lys
 85 90 95
 Gln Ile Phe Leu Gly Gly Val Asp Lys His Thr Gln Phe Trp Arg Tyr
 100 105 110
 Phe Ala Gly Asn Leu Ala Ser Gly Gly Ala Ala Gly Ala Thr Ser Leu
 115 120 125
 Lys Ile Val Tyr Pro Leu Asp Phe Ala Arg Thr Arg Leu Ala Ala Asp
 130 135 140
 Val Gly Lys Ser Gly Thr Glu Arg Glu Phe Arg Gly Leu Gly Asp Cys
 145 150 155 160
 Leu Val Lys Ile Thr Lys Ser Asp Gly Ile Arg Gly Leu Tyr Gln Gly
 165 170 175
 Phe Ser Val Ser Val Gln Gly Ile Ile Ile Tyr Arg Ala Ala Tyr Ile
 180 185 190
 Gly Val Tyr Asp Thr Ala Lys Gly Met Leu Thr Asp Pro Lys Asn Thr
 195 200 205
 His Ile Val Val Ser Trp Met Ile Ala Gln Thr Val Thr Ala Val Ala
 210 215 220

